

MLLNVLRICI	IVCLVNDGAG	KHSEGRERTK	TYSLNSRGYF	40
RKERGARRSK	ILLVNTKGLD	EPHIGHGDFG	LVAELFDSTR	80
THTNRKEPDM	NKVKLFSTVA	HG <u>NKS</u> ARRKA	<u>YNGS</u> RRNIFS	120
RRSFDKRNTE	VTEKPGAKMF	WNNFLVKMNG	APQNT <u>SH</u> GSK	160
AQEIMKEACK	TLPFTQNIVH	ENCDRMVION	NLCFGKCISL	200
HVPNQQDRRN	TCSHCLPSKF	TLNHLTL <u>NCT</u>	GSKNVVKVVM	240
MVEECTCEAH	KSNFHQTAQF	NMDTSTTLHH		270

Figure 1. Deduced amino acid sequence of *Xenopus cerberus* protein. SEQ ID NO:1.

Figure 2. Nucleotide sequence of the full-length cerberus DNA derived from the *Xenopus* organizer. The sense strand is on top (in the 5' to 3' direction) and the antisense strand on the bottom line (on the opposite direction). SEQ ID NO:2.

GAATTC	CCAG	TCGCTC	AGAA	CACTG	CAGG	GTCTAG	ATAT	CATACA	ATGT	TACTAA	60	
CCTA	AGGG	GTTC	AGCG	AG	TCTT	TGTGAC	GTCC	CAGATC	TATAG	TATGT	TACA	ATGATT
ATGT	ACTCAG	GATCT	GTATT	ATCG	TCTGCC	TTGT	GAATGA	TGGAG	CAGGA	AAAC	ACTCAG	120
TACAT	GAGTC	CTAG	ACATAA	TAGC	AGACGG	AACAC	TTACT	ACCT	CGTCCT	TTTG	TGAGTC	
AAGG	ACGAGA	AAGG	ACAAAA	ACAT	ATTAC	TTAAC	AGCAG	AGGT	TACTTC	AGAAA	AAGAAA	180
TTCT	TGCTCT	TTCT	TGTTTT	TGTAT	AAGTG	AATT	TGCGTC	TCCA	ATGAAG	TCTTT	TCTTT	
GAGG	AGCACG	TAGG	AGCAAG	ATTCT	GTCTGG	TGAAT	ACTAA	AGGT	CTTGAT	GAAC	CCCCACA	240
CTCT	CGTGC	ATCT	CGTTC	TAAG	ACGACC	ACTT	ATGATT	TCCAG	AACTA	CTTG	GGGGTGT	
TTGG	GCATGG	TGAT	TTTCGC	TTAG	TAGCTG	AACT	ATTTGA	TTCC	ACCAGA	ACAC	ATACAA	300
AACCC	GTACC	ACTAA	AAGCG	AATCAT	CGAC	TTGATA	AACT	AAGT	TGGTCT	TGTG	TATGTT	
ACAG	AAAAGA	GCCAG	ACATG	AACAA	AGTCA	AGCT	TTTCTC	AACAG	TTGCC	CATG	GAAACA	360
TGTCT	TTTCT	CGGT	CTGTAC	TTGT	TTTCA	TCGAA	AAGAG	TTGT	CAACGG	GTAC	CTTTGT	
AAAG	TGCAAG	AAGAA	AGCT	TACAT	TGGT	CTAGA	AGGAA	TATTT	TTCT	CGCC	GTCTTT	420
TTTCA	CGTTC	TTCT	TTTTCGA	ATGT	TACCA	GATCT	TCCTT	ATAAA	AAGGA	GCGG	CAAGAA	
TTGA	TAAAG	AAATAC	AGAG	GTTACT	TGAA	AGCT	TGGTGC	CAAG	ATGTT	TGGA	ACAATT	480
AACT	ATTTTC	TTTAT	GTCTC	CAAT	GACTTT	TCGG	ACCACG	GTCT	TACAAG	ACCT	TGTAA	
TTTT	GGTTAA	AATGA	TGGA	GCCCC	ACAGA	ATACA	AGCCA	TGGC	AGTAA	GCAC	AGGAAA	540
AAAAC	CAATT	TTACT	TACCT	CGGG	TGTCT	TATGT	TCGGT	ACCG	TCAATT	CGTG	TCCTTT	
TAAT	GAAAGA	AGCT	TGCAAA	ACCT	TGTTTT	TCACT	CAGAA	TATT	TGACAT	GAAA	ACTGTG	600
ATTAC	TTTCT	TCGA	ACGTTT	TGGA	ACAAAA	AGTG	AGTCTT	ATAA	CATGTA	CTTT	TGACAC	
ACAG	GATGGT	GATAC	AGAAC	AATCT	TGTGCT	TTGGT	AAATG	CATCT	CTCTC	CATG	TTCCAA	660
TGTC	TACCA	CTAT	GTCTTG	TTAG	ACACGA	AACCA	TTTAC	GTAG	AGAGAC	GTACA	AGGTT	
ATCAG	CAAGA	TCGAC	GAAAT	ACTT	GTTC	ATTG	CTTGCC	GTCC	AAATTT	ACC	TGAACC	720
TAGT	CGTTCT	AGCT	GCTTTA	TGA	ACAAGG	TAAC	GAACGG	CAGG	TTTAAA	TGGG	ACTTGG	
ACCT	GACGCT	GAATT	GTACT	GGAT	CTAAGA	ATGT	AGTAAA	GGTT	GTCA	TG	GTAGAGG	780
TGG	ACTGCGA	CTTA	ACATGA	CCTA	GATTCT	TACAT	CATTT	CCA	ACAGTAC	TACC	ATCTCC	
AATG	CACGTG	TGAAG	CTCAT	AAGAG	CAACT	TCCAC	CAAAC	TGC	ACGTTT	AAC	ATGGATA	840
TTAC	GTGCAC	ACTTC	GAGTA	TTCT	CGTTGA	AGGT	GTTTG	ACGT	GTCAAA	TTGT	ACCTAT	
CATC	TACTAC	CCTGC	ACCAT	TAAAG	GA	CTG	AGT	CCAT	ACAGTA	TGGAA	ATGCC	900
GTAG	ATGATG	GGAC	GTGTA	ATTT	CCTGAC	GGT	ATGTCAT	ACCT	TTACGG	GAAA	ACAACC	
AAT	ATTTGTT	ACATA	CTATG	CATCT	AAAGC	ATTAT	GTTGC	CTT	CTATTT	ATATA	ACCAC	960
TTATA	AAACAA	TGAT	GATAC	GTAG	ATTTG	TAATA	CAACG	GAAG	ATAAG	TATAT	TGGTG	
ATG	GAATAAG	GATT	GTATGA	ATTATA	ATTA	ACAA	TGGCA	TTTT	GTGTAA	CATG	CAAGAT	1020
TAC	CTTATTC	CTAAC	ATACT	TAAT	ATTAAT	TGTT	TACCGT	AAAA	CACATT	GTAC	GTCTTA	

CTCTGTTCCA TCAGTTGCAA GATAAAAGGC AATATTTGTT TGACTTTTTT TCTACAAAAT 1080
 GAGACAAGGT AGTCAACGTT CTATTTTCCG TTATAAACAA ACTGAAAAAA AGATGTTTTA
 GAATACCCAA ATATATGATA AGATAATGGG GTCAAACTG TTAAGGGGTA ATGTAATAAT 1140
 CTTATGGGTT TATATACTAT TCTATTACCC CAGTTTGGAC AATCCCCAT TACATTATTA
 AGGGACTAAG TTTGCCCAGG AGCAGTGACC CATAACAACC AATCAGCAGG TATGATTAC 1200
 TCCCTGATTC AAACGGGTCC TCGTCACTGG GTATTGTTGG TTAGTCGTCC ATACTAAATG
 TGGTCACCTG TTTAAAAGCA AACATCTTAT TGGTTGCTAT GGGTTACTGC TTCTGGGCAA 1260
 ACCAGTGGAC AAATTTTCGT TTGTAGAATA ACCAACGATA CCCAATGACG AAGACCCGTT
 AATGTGTGCC TCATAGGGGG GTTAGTGTGT TGTGTACTGA ATAAATTGTA TTTATTTTCA 1320
 TTACACACGG AGTATCCCC CAATCACACA ACACATGACT TATTTAACAT AAATAAAGTA
 TGTTACAAAA AAAAAAAA
 ACAATGTTTT TTTTTTTT

Fig. 2. (Continuation page 2, SEQ ID NO:2).

MSRTRKVDL LLLAIPGLAL LLLPNAYCAS CEPVRIPMCK SMPWNMTKMP NHLHHSTQAN 60
AILAIEQFEG LLTTECSQDL LFFLCAMYAP ICTIDFQHEP IKPCKSV CER ARAGCEPILI 120
KYRHTWPESL ACEELPVYDR GVCISPEAIV TVEQGTDSMP DFSMDSNNGN CGSGREHCKC 180
KPMKATQKTY LKNNYNYVIR AKVKEVKVVC HDATAIVEVK EILKSSLVNI PKDTVTLYTN 240
SGCLCPQLVA NEEYIIMGYE DKERTLLLV EGSLAEKWRD RLAKVKRWD QKLRRPRKSK 300
DPVAPIPNKN SNSRQARS

Figure 3. Deduced amino acid sequence of *Xenopus* frazzled protein. SEQ ID NO:3.

Figure 4. Nucleotide sequence of the full-length frazzled cDNA derived from the *Xenopus* organizer. The sense strand of the DNA on top (5' to 3' direction) and the antisense strand on the bottom line (opposite direction). SEQ ID NO:4.

GAATTCCTT TCACACAGGA CTCTGGCAG AGGTGAATGG TTAGCCCTAT GGATTGGTT	60
CTTAAGGGAA AGTGTGTCT GAGGACCGTC TCCACTTACC AATCGGGATA CCTAAACCAA	
TGTTGATTTT GACACATGAT TGATTGCTTT CAGATAGGAT TGAAGGACTT GGATTTTAT	120
ACAACTAAAA CTGTGTAATA ACTAACGAAA GTCTATCCTA ACTTCCTGAA CCTAAAAATA	
CTAATTCGTC ACTTTTAAAT TATCTGAGTA ATTGTTTATT TTGTATTGGA TGGGACTAAA	180
GATTAAGACG TGAAAATTTA ATAGACTCAT TAACAAGTAA AACATAACCT ACCCTGATTT	
GATAAACTTA ACTCCTTGCT TTTGACTTGC CCATAAACTA TAAGGTGGGG TGAGTTGTAG	240
CTATTTGAAT TGAGGAACGA AACTGAACG GGTATTTGAT ATTCCACCCC ACTCAACATC	
TTGCTTTTAC ATGTGCCCAG ATTTTCCCTG TATTCCCTGT ATTCCCTCTA AAGTAAGCCT	300
AACGAAATG TACACGGGTC TAAAGGGAC ATAAGGGACA TAAGGGAGAT TTCATTCCGA	
ACACATACAG GTTGGGCAGA ATAACAATGT CTCGAACAAG GAAAGTGGAC TCATTACTGC	360
TGTGTATGTC CAACCCGTCT TATTGTTACA GAGCTTGTTT CTTTCACCTG AGTAATGACG	
TACTGGCCAT ACCTGGACTG GCGCTTCTCT TATTACCCAA TGCTTACTGT GCTTCGTGTG	420
ATGACCGGTA TGGACCTGAC CGCGAAGAGA ATAATGGGTT ACGAATGACA CGAAGCACAC	
AGCCTGTGCG GATCCCCATG TGCAAACTTA TGCCATGGAA CATGACCAAG ATGCCCAACC	480
TCGGACACGC CTAGGGGTAC ACGTTTAGAT ACGGTACCTT GTACTGGTTC TACGGGTTGG	
ATCTCCACCA CAGCACTCAA GCCAATGCCA TCCTGGCAAT TGAACAGTTT GAAGGTTTGC	540
TAGAGGTGGT GTCGTGAGTT CGGTTACGGT AGGACCGTTA ACTTGTCAAA CTTCCAAACG	
TGACCACTGA ATGTAGCCAG GACCTTTTGT TCTTCTGTG TGCCATGTAT GCCCCATTT	600
ACTGGTGACT TACATCGGTC CTGGAAAACA AGAAGACAC ACGGTACATA CGGGGGTAAA	
GTACCATCGA TTCCAGCAT GAACCAATTA AGCCTTGCAA GTCCGTGTGC GAAAGGGCCA	660
CATGGTAGCT AAAGGTCGTA CTTGGTTAAT TCGGAACGTT CAGGCACACG CTTTCCCGGT	
GGGCCGGCTG TGAGCCCATC CTCATAAAGT ACCGGCACAC TTGGCCAGAG AGCCTGGCAT	720
CCCGGCCGAC ACTCGGGTAA GAGTATTTC TGGCCGTGTG AACCGGTCTC TCGGACCGTA	
GTGAAGAGCT GCCCGTATAT GACAGAGGAG TCTGCATCTC CCCAGAGGCT ATCGTCACAG	780
CACCTCTCGA CGGGCATATA CTGTCTCCTC AGACGTAGAG GGGTCTCCGA TAGCAGTGTG	
TGGAACAAGG AACAGATTCA ATGCCAGACT TCTCCATGGA TTCAAACAAT GGAAATTGCG	840
ACCTTGTTCC TTGTCTAAGT TACGGTCTGA AGAGGTACCT AAGTTTGTTA CCTTTAACGC	
GAAGCGGCAG GGAGCACTGT AAATGCAAGC CCATGAAGGC AACCCAAAAG ACGTATCTCA	900
CTTCGCCGTC CCTCGTGACA TTTACGTTCC GGTACTTCCG TTGGGTTTTT TGCATAGAGT	
AGAATAATTA CAATTATGTA ATCAGAGCAA AAGTGAAAGA GGTGAAAGTG AAATGCCAGG	960
TCTTATTAAT GTTAATACAT TAGTCTCGTT TTCACTTTCT CCACTTTCAC TTTACGGTGC	
ACGCAACAGC AATTGTGGAA GTAAAGGAGA TTCTCAAGTC TTCCCTAGTG AACATTCTTA	1020
TGCGTTGTCT TTAACACCTT CATTTCTCT AAGAGTTTCA AAGGGATCAC TTGTAAGGAT	

AAGACACAGT GACACTGTAC ACCAACTCAG GCTGCTTGTG CCCCCAGCTT GTTGCCAATG TTCTGTGTCA CTGTGACATG TGGTTGAGTC CGACGAACAC GGGGGTCGAA CAACGGTTAC	1080
AGGAATACAT AATTATGGGC TATGAAGACA AAGAGCGTAC CAGGCTTCTA CTAGTGGAAAG TCCTTATGTA TTAATACCCG ATACTTCTGT TTCTCGCATG GTCCGAAGAT GATCACCTTC	1140
GATCCTTGGC CGAAAAATGG AGAGATCGTC TTGCTAAGAA AGTCAAGCGC TGGGATCAAA CTAGGAACCG GCTTTTTTACC TCTCTAGCAG AACGATTCTT TCAGTTCGCG ACCCTAGTTT	1200
AGCTTCGACG TCCCAGGAAA AGCAAAGACC CCGTGGGTCC AATTCCCAAC AAAACAGCA TCGAAGCTGC AGGGTCCTTT TCGTTTCTGG GGCACCGAGG TTAAGGGTTG TTTTGTCTGT	1260
ATTCCAGACA AGCGCGTAGT TAGACTAACG GAAAGGTGTA TGGAACTCT ATGGACTTTG TAAGGTCTGT TCGCGCATCA ATCTGATTGC CTTCCACAT ACCTTTGAGA TACCTGAAAC	1320
AAACTAAGAT TTGCATTGTT GGAAGAGCAA AAAAGAAATT GCACTACAGC ACGTTATATT TTTGATTCTA AACGTAACAA CCTTCTCGTT TTTTCTTTAA CGTGATGTCG TGCAATATAA	1380
CTATTGTTTA CTACAAGAAG CTGGTTTAGT TGATTGTAGT TCTCCTTTCC TTCTTTTTTT GATAACAAAT GATGTTCTTC GACCAAATCA ACTAACATCA AGAGGAAAGG AAGAAAAAAA	1440
TTATAACTAT ATTTGCACGT GTTCCCAGGC AATGTTTTTA TTCAACTTCC AGTGACAGAG AATATTGATA TAAACGTGCA CAAGGGTCCG TTAACAAAT AAGTTGAAGG TCACTGTCTC	1500
CAGTGACTGA ATGTCTCAGC CTAAAGAAGC TCAATTCATT TCTGATCAAC TAATGGTGAC GTCAGTACT TACAGAGTCG GATTTCTTCG AGTTAAGTAA AGACTAGTTG ATTACCACTG	1560
AAGTGTGTTGA TACTTGGGGA AAGTGAACATA ATTGCAATGG TAAATCAGAG AAAAGTTGAC TTCACAACT ATGAACCCCT TTCACTTGAT TAACGTTACC ATTTAGTCTC TTTTCAACTG	1620
CAATGTTGCT TTTCTGTAG ATGAACAAGT GAGAGATCAC ATTTAAATGA TGATCACTTT GTTACAACGA AAAGGACATC TACTTGTTCA CTCTCTAGTG TAAATTTACT ACTAGTGAAA	1680
CCATTTAATA CTTTCAGCAG TTTTAGTTAG ATGACATGTA GGATGCACCT AAATCTAAAT GGTAAATTAT GAAAGTCGTC AAAATCAATC TACTGTACAT CCTACGTGGA TTTAGATTTA	1740
ATTTTATCAT AAATGAAGAG CTGGTTTAGA CTGTATGGTC ACTGTTGGGA AGGTAAATGC TAAATAGTA TTTACTTCTC GACCAAATCT GACATACCAG TGACAACCCCT TCCATTACG	1800
CTACTTTGTC AATTCTGTTT TAAAAATTGC CTAAATAAAT ATTAAGTCCT AAATAAAAAA GATGAAACAG TTAAGACAAA ATTTTAAACG GATTATTTA TAATTCAGGA TTTATTTTTT	1860
AAAAAAAAAA AAAAA TTTTTTTTTT TTTT	

Fig. 4. (Continuation page 2, SEQ ID NO:4).

MLLLFRAIPM LLLGLMVLQT DCEIAQYYID EEEPPGTVIA VLSQHSIFNT TDIPATNFRL	60
MKQFNNSLIG VRESQGQLSI MERIDREQIC RQSLHCNLAL DVVSFSKGHF KLLNVKVEVR	120
DINDHSPHFP SEIMHVEVSE SSSVGTRIPL EIAIDEDVGS NSIQNFQISN NSHFSIDVLT	180
RADGVKYADL VLMRELDREI QPTYIMELLA MDGGVPSLSG TAVVNIRVLD FNDNSPVFER	240
STIAVDLVED APLGYLLEL HATDDDEGVN GEIVYGFSTL ASQEVRLFK INSRTGSVTL	300
EGQVDFETKQ TYEFEVQAQD LGPNPLTATC KVTVHILDVN DNTPAITITP LTTVNAGVAY	360
IPETATKENF IALISTTDRA SGSNGQVRCT LYGHEHFKLQ QAYEDSYMIV TTSTLDRENI	420
AAYSILTVAE DLGFPPLKTK KYITVKVSE NDNAPVFSKP QYEASILENN APGSYITTVI	480
ARDSDSQNG KVNRLVDAK VMQSLTTFV SLDADSGVLR AVRSLDYEKL KQLDFEIEAA	540
DNGIPQLSTR VQLNLRIVDQ NDNCPVITNP LLNNGSGEVL LPISAPQNYL VFQLKAEDSD	600
EGHNSQLFYT ILRDPSRLFA INKESGEVFL KKQLNSDHSE DLSIVVAVYD LGRPSLSTNA	660
TVKFILTDSF PSNVEVVILQ PSAEEQHQID MSIIFIIVLA GGCALLLLAI FFVACTCKKK	720
AGEFKQVPEQ HGTCNEERLL STPSQSVSS SLSQSESCQL SINTESENCV VSSNQEQHQQ	780
TGIKHSISVP SYHTSGWHLN NCAMSSISGHS HMGHISTKVQ WAKEIVTSMT VTLILVENQK	840
RRALSSQCRH KPVLTQMNQ QGSDMPITIS ATESTRVQKM GTAHCMKRA IDCLTL	

Figure 5. Deduced amino acid sequence of the *Xenopus* PAPC (paraxial protocadherin) protein. It encodes a member of the cadherin family of transmembrane proteins that has dorsalizing activity when constructs are injected into *Xenopus* embryos. SEQ ID NO:5.

Figure 6. Nucleotide sequence of the full-length PAPC cDNA derived from the *Xenopus* organizer. The sense strand of the DNA is shown in the top line (in the 5' to 3' direction), and the bottom line shows the antisense strand (opposite orientation). SEQ ID NO:6.

GAATTC	CCAG	AGATGA	ACTC	CTTGAG	ATTG	TTTTAA	TGA	CTGCAG	GTCT	GGAAG	GATT	60
CTTAAG	GGTC	TCTACT	TGAG	GAAC	TCTAAC	AAAATT	TACT	GACG	TCCAGA	CCTTC	CTAAG	
ACATTG	CCAC	ACTGTT	TCTA	GGCAT	GAAAA	AAC	TGCAAGT	TTCA	ACTTTG	TTTTG	GTGC	120
TGTAAC	GGTG	TGACAA	AGAT	CCGTACT	TTTT	TTGAC	GTTC	AAGTT	GAAAC	AAAA	ACCAG	
AAC	TTTGATT	CTTCA	AGATG	CTGCTT	CTCT	TCAGAG	CCAT	TCCA	ATGCTG	CTGTT	GGGAC	180
TTGAA	ACTAA	GAAGT	TCTAC	GACGA	AGAGA	AGTCT	CGGT	AGGT	TACGAC	GACA	ACCTG	
TGATG	GGTTTT	ACAA	ACAGAC	TGTGA	AATTG	CCCAG	TACTA	CATAG	ATGAA	GAAGA	ACCCC	240
ACTAC	CAAAA	TGTTT	GTCTG	ACACTT	TAAC	GGGTC	CATGAT	GTATC	TACTT	CTTCT	TGGGG	
CTGGC	ACTGT	AATTG	CAGTG	TTGTC	CACAAC	ACTCC	ATATT	TAACA	CTACA	GATAT	ACCTG	300
GACCG	TGACA	TTAAC	GTAC	AACAG	TGTTG	TGAGG	TATAA	ATTGT	GATGT	CTAT	TGGAC	
CAACCA	ATTT	CCGTCT	AAATG	AAGCA	ATTTA	ATAAT	TCCCT	TATCG	GAGTC	CGTG	GAGTG	360
GTTGG	TAAA	GGCAG	ATTAC	TTCGT	TAAAT	TATTA	AGGGA	ATAGC	CTCAG	GCACT	CTCAC	
ATGGG	CAGCT	GAGCAT	CATG	GAGAG	GATTG	ACCGG	GAGCA	AATCT	GCAGG	CAGTC	CCCTTC	420
TACCC	GTCGA	CTCGT	AGTAC	CTCTC	CTAAC	TGGCC	CTGT	TTAGA	CGTCC	GTCAG	GGAAG	
ACTG	CAACCT	GGCTT	TGGAT	GTGGT	CAGCT	TTTCC	AAAGG	ACACT	TCAAG	CTTCT	GAAAG	480
TGACG	TTGGA	CCGAA	ACCTA	CACCAG	TGCA	AAAGG	TTTCC	TGTGA	AGTTC	GAAGA	CTGC	
TGAA	AGTGA	GGTG	AGAGAC	ATTA	ATGACC	ATAGC	CCCTCA	CTTTC	CCAGT	GAAAT	AATGC	540
ACTTT	CACCT	CCACT	CTCTG	TAATT	ACTGG	TATCG	GAGAGT	GAAAG	GGTCA	CTTT	TATTAG	
ATGT	GAGGT	GTCTG	AAAGT	TCCTC	TGTGG	GCACC	AGGAT	TCCTT	TAGAA	ATTG	CAATAG	600
TACAC	CTCA	CAGAC	TTTCA	AGGAG	ACACC	CGTGG	TCCTA	AGGAA	ATCTT	TAACG	TTATC	
ATGA	AGATGT	TGGGT	CCAAC	TCCAT	CCAGA	ACTTT	CAGAT	CTCAA	ATAAT	AGCCA	CTTCA	660
TACTT	CTACA	ACCCAG	GTG	AGGTAG	GTCT	TGAA	AGTCTA	GAGTT	TATTA	TCGGT	GAAGT	
GCATT	GATGT	GCTAA	ACCAGA	GCAGAT	GGGG	TGAA	ATATGC	AGATT	TAGTC	TTAAT	GAGAG	720
CGTAA	CTACA	CGATT	GGTCT	CGTCT	ACCCC	ACTTT	TATAG	TCTAA	ATCAG	AATT	ACTCTC	
AACTG	GACAG	GGAA	ATCCAG	CCAAC	ATACA	TAATG	GAGCT	ACTAG	CAATG	GATGG	GGGTG	780
TTGAC	CTGC	CCTTT	AGGTC	GGTTG	TATGT	ATTAC	CTCGA	TGATC	GTTC	CTAC	CCCCAC	
TACCA	CTACT	ATCTG	GTACT	GCAGT	GGTTA	ACATC	CGAGT	CCTGG	ACTTT	AATGA	TAAACA	840
ATGGT	AGTGA	TAGAC	CATGA	CGTCA	CCAAT	TGTAG	GCTCA	GGACC	TGAAA	TTACT	TATTGT	
GCCC	AGTGT	TGAGA	GAAGC	ACCAT	TGCTG	TGGAC	CTAGT	AGAGG	ATGCT	CCTCT	GGGAT	900
CGGGT	CACAA	ACTCT	CTTCG	TGGTA	ACGAC	ACCTG	GATCA	TCTCT	ACGA	GGAG	ACCCTA	
ACCTT	TTTGT	GGAGT	TACAT	GCTAC	TGACG	ATGAT	GAAGG	AGTGA	ATGGA	GAAAT	TGTTT	960
TGGA	AAACAA	CCTCA	ATGTA	CGATG	ACTGC	TACTA	CTTCC	TCACT	TACCT	CTTTA	ACAAA	
ATGG	ATTAG	CCTTT	TGGCA	TCTCA	AGAGG	TACGT	CAGCT	ATTTA	AAAT	AATCC	AGAA	1020
TACCT	AAGTC	GTGA	AAACGT	AGAGT	TCTCC	ATGCA	GTCGA	TAAAT	TTTTAA	TTGAG	GTCTT	

CTGGCAGTGT TACTCTTGAA GGCCAAGTTG ATTTTGAGAC CAAGCAGACT TACGAATTTG 1080
 GACCGTCACA ATGAGAACTT CCGGTTCAAC TAAAACTCTG GTTCGTCTGA ATGCTTAAAC
 AGGTACAAGC CCAAGATTG GGCCCCAACC CACTGACTGC TACTTGTAAG GTAACGTGTC 1140
 TCCATGTTCC GGTCTAAAC CCGGGGTGG GTGACTGACG ATGAACATTT CATTGACAAG
 ATATACTTGA TGTAATGAT AATACCCAG CCATCACTAT TACCCCTCTG ACTACTGTAA 1200
 TATATGAAC ACATTTACTA TTATGGGTC GGTAGTGATA ATGGGAGAC TGATGACATT
 ATGCAGGAGT TGCCTATATT CCAGAAACAG CCACAAAGGA GAACTTTATA GCTCTGATCA 1260
 TACGTCCTCA ACGGATATAA GGTCTTTGTC GGTGTTTCCT CTTGAAATAT CGAGACTAGT
 GCACTACTGA CAGAGCCTCT GGATCTAATG GACAAGTTCG CTGTACTCTT TATGGACATG 1320
 CGTGATGACT GTCTCGGAGA CCTAGATTAC CTGTTCAAGC GACATGAGAA ATACCTGTAC
 AGCACTTTAA ACTACAGCAA GCTTATGAGG ACAGTTACAT GATAGTTACC ACCTCTACTT 1380
 TCGTGAAATT TGATGTCGTT CGAATACTCC TGCAATGTA CTATCAATGG TGGAGATGAA
 TAGACAGGGA AAACATAGCA GCGTACTCTT TGACAGTAGT TGCAGAAGAC CTTGGCTTCC 1440
 ATCTGTCCCT TTTGTATCGT CGCATGAGAA ACTGTCATCA ACGTCTTCTG GAACCGAAGG
 CCTCATTGAA GACCAAAAAG TACTACACAG TCAAGGTTAG TGATGAGAAT GACAATGCAC 1500
 GGAGTAACTT CTGGTTTTTC ATGATGTGTC AGTTCCAATC ACTACTCTTA CTGTTACGTG
 CTGTATTTTC TAAACCCAG TATGAAGCTT CTATTCTGGA AAATAATGCT CCAGGCTCTT 1560
 GACATAAAG ATTTGGGGTC ATACTTCGAA GATAAGACCT TTTATTACGA GGTCCGAGAA
 ATATAACTAC AGTGATAGCC AGAGACTCTG ATAGTGATCA AAATGGCAA GTAAATTACA 1620
 TATATTGATG TCACTATCGG TCTCTGAGAC TATCACTAGT TTTACCGTTT CATTTAATGT
 GACTTGTGGA TGCAAAAGTG ATGGGCCAGT CACTAACAAC ATTTGTTTCT CTTGATGCGG 1680
 CTGAACACCT ACGTTTTAC TACCCGGTCA GTGATTGTTG TAAACAAAGA GAACTACGCC
 ACTCTGGAGT ATTGAGAGCT GTTAGGTCTT TAGACTATGA AAACTTAA CAACTGGATT 1740
 TGAGACCTCA TAACTCTCGA CAATCCAGAA ATCTGATACT TTTTGAATT GTTGACCTAA
 TTGAAATTGA AGCTGCAGAC AATGGGATCC CTCAACTCTC CACTCGCGTT CAACTAAATC 1800
 AACTTTAACT TCGACGCTG TTACCCTAGG GAGTTGAGAG GTGAGCGCAA GTTGATTAG
 TCAGAATAGT TGATCAAAAT GATAATTGCC CTGTGATAAC TAATCCTCTT CTTAATAATG 1860
 AGTCTTATCA ACTAGTTTAA CTATTAACGG GACACTATTG ATTAGGAGAA GAATTATTAC
 GCTCGGGTGA AGTTCTGCTT CCCATCAGCG CTCCTCAAAA CTATTTAGTT TTCCAGCTCA 1920
 CGAGCCCACT TCAAGACGAA GGGTAGTCGC GAGGAGTTT GATAAATCAA AAGGTCGAGT
 AAGCCGAGGA TTCAGATGAA GGGCACAAC CCCAGCTGTT CTATACCATA CTGAGAGATC 1980
 TTCGGCTCCT AAGTCTACTT CCCGTGTTGA GGGTCGACAA GATATGGTAT GACTCTCTAG
 CAAGCAGATT GTTTGCCATT AACAAAGAAA GTGGTGAAGT GTTCCTGAAA AAACAATTAA 2040
 GTTCGTCTAA CAAACGGTAA TTGTTTCTTT CACCCTTCA CAAGGACTTT TTTGTTAATT
 ACTCTGACCA TTCAGAGGAC TTGAGCATAG TAGTTGCAGT GTATGACTTG GGAAGACCTT 2100
 TGAGACTGGT AAGTCTCTG AACTCGTATC ATCAACGTCA CATACTGAAC CCTTCTGGAA
 CATTATCCAC CAATGCTACA GTTAAATTCA TCCTCACCAG CTCTTTTCCT TCTAACGTTG 2160
 GTAATAGGTG GTTACGATGT CAATTTAAGT AGGAGTGGCT GAGAAAAGGA AGATTGCAAC

Fig. 6. (Continuation page 2, SEQ ID NO:6).

AAGTCGTTAT TTTGCAACCA TCTGCAGAAG AGCAGCACCA GATCGATATG TCCATTATAT 2220
TTCAGCAATA AAACGTTGGT AGACGTCTTC TCGTCGTGGT CTAGCTATAC AGGTAATATA
TCATTGCAGT GCTGGCTGGT GGTGTGTGCTT TGCTACTTTT GGCCATCTTT TTTGTGGCCT 2280
AGTAACGTCA CGACCGACCA CCAACACGAA ACGATGAAAA CCGGTAGAAA AAACACCGGA
GTACTTGTA AAAGAAAGCT GGTGAATTTA AGCAGGTACC TGAACAACAC GGAACATGCA 2340
CATGAACATT TTTCTTTTGA CCACTTAAAT TCGTCCATGG ACTTGTGTG CCTGTACGT
ATGAAGAACG CCTGTTAAGC ACCCATCTC CCCAGTCGGT CTCTTCTTCT TTGTCTCAGT 2400
TACTTCTTGC GGACAATTCG TGGGGTAGAG GGGTCAGCCA GAGAAGAAGA AACAGAGTCA
CTGAGTCATG CCAACTCTCC ATCAATACTG AATCTGAGAA TTGCAGCGTG TCCTCTAACC 2460
GACTCAGTAC GGTGAGAGG TAGTTATGAC TTAGACTCTT AACGTCGCAC AGGAGATTGG
AAGAGCAGCA TCAGCAAACA GGCATAAAGC ACTCCATCTC TGTACCATCT TATCACACAT 2520
TTCTCGTCGT AGTCGTTTGT CCGTATTTTG TGAGGTAGAG ACATGGTAGA ATAGTGTGTA
CTGGTTGGCA CCTGGACAAT TGTGCAATGA GCATAAGTGG ACATTCTCAC ATGGGGCACA 2580
GACCAACCGT GGACCTGTTA ACACGTTACT CGTATTCACC TGTAAGAGTG TACCCCGTGT
TTAGTACAAA GGTACAGTGG GCAAAGGAGA TAGTGACTTC AATGACAGTG ACTCTGATAC 2640
AATCATGTTT CCATGTCACC CGTTTCTCT ATCACTGAAG TTACTGTCAC TGAGACTATG
TAGTGGAGAA TCAGAAAAGA AGAGCATTGA GCAGCCAATG CAGGCACAAG CCAGTGCTCA 2700
ATCACCTCTT AGTCTTTTCT TCTCGTAACT CGTCGGTTAC GTCGGTGTTC GGTCACGAGT
ATACACAGAT GAATCAGCAG GGTTCGACA TGCCGATAAC TATTCAGCC ACCGAATCAA 2760
TATGTGTCTA CTTAGTCGTC CCAAGGCTGT ACGGCTATTG ATAAAGTCGG TGGCTTAGTT
CAAGGGTCCA GAAATGGGA ACTGCACATT GCAATATGAA AAGGGCTATA GACTGTCTTA 2820
GTTCCAGGT CTTTACCCT TGACGTGTAA CGTTATACTT TTCCCGATAT CTGACAGAAT
CTCTGTAGCT CCTGTATATT ACAATACCTA CCATGCAAGA ATGCCTAACC TGCACATACC 2880
GAGACATCGA GGACATATAA TGTTATGGAT GGTACGTTCT TACGGATTGG ACGTGTATGG
GAACCATACC CTTAGAGACC CTTATTACCA TATCAATAAT CCTGTTGCTA ATCGGATGCA 2940
CTTGGTATGG GAATCTCTGG GAATAATGGT ATAGTTATTA GGACAACGAT TAGCCTACGT
GGCGGAATAT GAAAGAGATT TAGTCAACAG AAGTGAACG TTATCTCCGC AGAGATCGTC 3000
CCGCTTATA CTTTCTCTAA ATCAGTTGTC TTCAGTTGC AATAGAGGCG TCTCTAGCAG
TAGCAGATAC CAAGAATTCA ATTACAGTCC GCAGATATCA AGACAGCTTC ATCCTTCAGA 3060
ATCGTCTATG GTTCTTAAGT TAATGTCAGG CGTCTATAGT TCTGTGCAAG TAGGAAGTCT
AATTGCTACA ACCTTTTAAT CATTAGGCAT GCAAGTGAGA ATGCACAAAG GCAAGTGCTT 3120
TTAACGATGT TGGAAAATTA GTAATCCGTA CGTTCACTCT TACGTGTTTC CGTTCACGAA
TAGCATGAAA GCTAAATATA TGGAGTCTCC CCTTCCCTC TGATGGATGG GGGGAGACAC 3180
ATCGTACTTT CGATTTATAT ACCTCAGAGG GGAAAGGGAG ACTACCTACC CCCCTCTGTG
AGGACAGTGC ATAAATATAC AGCTGCTTTC TATTGCAAT TCACTTGGGA ATTTTGTGTT 3240
TCCTGTCACG TATTATATG TCGACGAAAG ATAAACGTAA AGTGAACCTT TAAAAACAA
TTTTTACAT ATTTATTTT CCTGAATTGA ATGTGACATT GTCCTGTCAC CTAAGTAGCA 3300
AAAAATGTA TAAATAAAAA GGACTTAACT TACACTGTAA CAGGACAGTG GATTGATCGT

Fig. 6. (Continuation page 3, SEQ ID NO:6).

ATTAAATCCA CAGACCTACA GTCAAATATT TGAGGGCCCC TGAAACAGCA CATCAGTCAG	3360
TAATTTAGGT GTCTGGATGT CAGTTTATAA ACTCCCGGGG ACTTTGTCGT GTAGTCAGTC	
GACCTAAAGT GGCCTTTTTA CTTTTCAGC CTCCTGGGTC TGCCCTCTGT GTTAATCAGC	3420
CTGGATTTC ACGGAAAAAT GAAAATCGTC GAGGACCCAG ACGGGAGACA CAATTAGTCG	
CCCTGGTCAA GTCCTGAGTA GGATCATGGC GTTTTATAT GCATCTCACC TACTTTGGAC	3480
GGGACCAGTT CAGGACTCAT CCTAGTACCG CAAAAATATA CGTAGAGTGG ATGAAACCTG	
GTGATTTACA CATAATAGGA AACGCTTGGT TTCAGTGAAG TCTGTGTTGT ATATATTCTG	3540
CACTAAATGT GTATTATCCT TTGCGAACCA AAGTCACTTC AGACACAACA TATATAAGAC	
TTATATACAC GCATTTTGTG TTTGTGTATA TATTTCAGT CCATTCAGAT ATGTGTATAT	3600
AATATATGTG CGTAAACAC AAACACATAT ATAAAGTTCA GGTAAGTCTA TACACATATA	
AGTGCAGACC TTGTAAATTA AATATTCTGA TACTTTTCC TCAATAAATA TTAAAT	
TCACGTCTGG AACATTTAAT TTATAAGACT ATGAAAAGG AGTTATTAT AAATTTA	

Fig. 6. (Continuation page 4, SEQ ID NO:6).

MVCCGPGRML LGWAGLLVLA ALCLLQVPGA QAAACEPVRI PLCKSLPWNM TKMPNHLHHS 60
TQANAILAME QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE 120
PILIKYRHSW PESLACDELP VYDRGVCISP EAIVTADGAD FPMDSSTGHC RGASSERCKC 180
KPV RATQKTY FRNNYNYVIR AKVKEVKMKC HDVTAVVEVK EILKASLVNI PRDTVNLYTT 240
SGCLCPPLTV NEEYVIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLGK 300
TDASDSTQNQ KSGRNSNPRP ARS.

Figure 7. Deduced amino acid sequence of mouse FRZB-1 protein. SEQ ID NO:7.

Figure 8. Nucleotide sequence of the full-length mouse FRZB-1 cDNA. SEQ ID NO:8.

AAGCCTGGGA CCATGGTCTG CTGCGGCCCG GGACGGATGC TGCTAGGATG GGCCGGGTTG	60
TTCCGACCCT GGTACCAGAC GACGCCGGGC CCTGCCTACG ACGATCCTAC CCGGCCCAAC	
CTAGTCCTGG CTGCTCTCTG CCTGCTCCAG GTGCCCGGAG CTCAGGCTGC AGCCTGTGAG	120
GATCAGGACC GACGAGAGAC GGACGAGGTC CACGGGCCTC GAGTCCGACG TCGGACACTC	
CCTGTCCGCA TCCCGCTGTG CAAGTCCCTT CCCTGGAACA TGACCAAGAT GCCCAACCAC	180
GGACAGGCGT AGGGCGACAC GTTCAGGGAA GGGACCTTGT ACTGGTTCTA CGGGTGTTGT	
CTGCACCACA GCACCCAGGC TAACGCCATC CTGGCCATGG AACAGTTCGA AGGGCTGCTG	240
GACGTGGTGT CGTGGGTCCG ATTGCGGTAG GACCGGTACC TTGTCAAGCT TCCCACGAC	
GGCACCCTACT GCAGCCCGGA TCTTCTCTTC TTCCTCTGTG CAATGTACGC ACCCATTTGC	300
CCGTGGGTGA CGTCGGGCTT AGAAGAGAAG AAGGAGACAC GTTACATGCG TGGGTAAACG	
ACCATCGACT TCCAGCACGA GCCCATCAAG CCCTGCAAGT CTGTGTGTGA GCGCGCCCGA	360
TGGTAGCTGA AGGTCGTGCT CGGGTAGTTC GGGACGTTCA GACACACACT CGCGCGGGCT	
CAGGGCTGCG AGCCCATTTCT CATCAAGTAC CGCCACTCGT GGCCGGAAAG CTTGGCCTGC	420
GTCCCAGCGC TCGGGTAAGA GTAGTTTCATG GCGGTGAGCA CCGGCCTTTC GAACCGGACG	
GACGAGCTGC CGGTGTACGA CCGCGGCGTG TGCATCTCTC CTGAGGCCAT CGTCACCGCG	480
CTGCTCGACG GCCACATGCT GGCGCCGCAC ACGTAGAGAG GACTCCGGTA GCAGTGGCGC	
GACGGAGCGG ATTTTCCTAT GGATTCAAGT ACTGGACACT GCAGAGGGGC AAGCAGCGAA	540
CTGCCTCGCC TAAAAGGATA CCTAAGTTCA TGACCTGTGA CGTCTCCCCG TTCGTCGCTT	
CGTTGCAAAT GTAAGCCTGT CAGAGCTACA CAGAAGACCT ATTTCCGGAA CAATTACAAC	600
GCAACGTTTA CATTCCGACA GTCTCGATGT GTCTTCTGGA TAAAGGCCTT GTTAATGTTG	
TATGTCATCC GGGCTAAAGT TAAAGAGGTA AAGATGAAAT GTCATGATGT GACCGCCGTT	660
ATAAGTAGG CCGGATTTC AATTCTCCAT TTCTACTTTA CAGTACTACA CTGGCGGCAA	
GTGGAAGTGA AGGAAATTCT AAAGGCATCA CTGGTAAACA TTCCAAGGGA CACCGTCAAT	720
CACCTTCACT TCCTTTAAGA TTCCGTAGT GACCATTGTG AAGGTTCCCT GTGGCAGTTA	
CTTTATACCA CCTCTGGCTG CCTCTGTCCT CCACTTACTG TCAATGAGGA ATATGTCATC	780
GAAATATGGT GGAGACCGAC GGAGACAGGA GGTGAATGAC AGTTACTCCT TATACAGTAG	
ATGGGCTATG AAGACGAGGA ACGTTCAGG TTACTCTTGG TAGAAGGCTC TATAGCTGAG	840
TACCCGATAC TTCTGCTCCT TGCAAGGTCC AATGAGAACC ATCTTCCGAG ATATCGACTC	
AAGTGGAAGG ATCGGCTTGG TAAGAAAGTC AAGCGCTGGG ATATGAAACT CCGACACCTT	900
TTCACCTTCC TAGCCGAACC ATTCTTTCAG TTCGCGACCC TATACTTTGA GGCTGTGGAA	
GGACTGGGTA AACTGATGC TAGCGATTCC ACTCAGAATC AGAAGTCTGG CAGGAAGTCT	960
CCTGACCCAT TTTGACTACG ATCGCTAAGG TGAGTCTTAG TCTTCAGACC GTCCTTGAGA	

AATCCCCGGC CAGCACGCAG CTAAATCCTG AAATGTAAAA GGCCACACCC ACGGACTCCC	1020
TTAGGGGGCCG GTCGTGCGTC GATTTAGGAC TTTACATTTT CCGGTGTGGG TGCCTGAGGG	
TTCTAAGACT GCGCTGGTG GACTAACAAA GGAAAACCGC ACAGTTGTGC TCGTGACCGA	1080
AAGATTCTGA CCGCGACCAC CTGATTGTTT CCTTTTGGCG TGTCAACACG AGCACTGGCT	
TTGTTTACCG CAGACACCGC GTGGCTACCG AAGTTACTTC CGGTCCCCTT TCTCCTGCTT	1140
AACAAATGGC GTCTGTGGCG CACCGATGGC TTCAATGAAG GCCAGGGGAA AGAGGACGAA	
CTTAATGGCG TGGGGTTAGA TCCTTTAATA TGTATATAT TCTGTTTCAT CAATCACGTG	1200
GAATTACCGC ACCCCAATCT AGGAAATTAT ACAATATATA AGACAAAGTA GTTAGTGCAC	
GGGACTGTTC TTTTGCAACC AGAATAGTAA ATTAAATATG TTGATGCTAA GGTTCCTGTA	1260
CCCTGACAAG AAAACGTTGG TCTTATCATT TAATTTATAC AACTACGATT CCAAAGACAT	
CTGGACTCCC TGGGTTTAAT TTGGTGTCT GTACCCTGAT TGAGAATGCA ATGTTTCATG	1320
GACCTGAGGG ACCCAAATTA AACCACAAGA CATGGGACTA ACTCTTACGT TACAAAGTAC	
TAAAGAGAGA ATCCTGGTCA TATCTCAAGA ACTAGATATT GCTGTAAGAC AGCCTCTGCT	1380
ATTTCTCTCT TAGGACCAGT ATAGAGTTCT TGATCTATAA CGACATTCTG TCGGAGACGA	
GCTGCGCTTA TAGTCTTGTG TTTGTATGCC TTTGTCCATT TCCCTCATGC TGTGAAAGTT	1440
CGACGCGAAT ATCAGAACAC AAACATACGG AAACAGGTAA AGGGAGTACG ACACTTTCAA	
ATACATGTTT ATAAAGGTAG AACGGCATTT TGAAATCAGA CACTGCACAA GCAGAGTAGC	1500
TATGTACAAA TATTTCCATC TTGCCGTAAA ACTTTAGTCT GTGACGTGTT CGTCTCATCG	
CCAACACCAG GAAGCATTTA TGAGGAAACG CCACACAGCA TGACTTATTT TCAAGATTGG	1560
GGTTGTGGTC CTTCGTAAAT ACTCCTTTGC GGTGTGTCGT ACTGAATAAA AGTTCTAACC	
CAGGCAGCAA AATAAATAGT GTTGGGAGCC AAGAAAAGAA TATTTTGCCT GGTTAAGGGG	1620
GTCCGTCGTT TTATTTATCA CAACCCTCGG TTCTTTTCTT ATAAAACGGA CCAATTCCCC	
CACACTGGAA TCAGTAGCCC TTGAGCCATT AACAGCAGTG TTCTTCTGGC AAGTTTTTGA	1680
GTGTGACCTT AGTCATCGGG AACTCGGTAA TTGTCGTCAC AAGAAGACCG TTCAAAAAC	
TTGTTTCATA AATGTATTCA CGAGCATTAG AGATGAACTT ATAAC TAGAC ATCTGTTGTT	1740
AAACAAGTAT TTACATAAGT GTCGTAATC TCTACTTGAA TATTGATCTG TAGACAACAA	
ATCTCTATAG CTCTGCTTCC TTCTAAATCA AACCCATTGT TGGATGCTCC CTCTCCATT	1800
TAGAGATATC GAGACGAAGG AAGATTTAGT TTGGGTAACA ACCTACGAGG GAGAGGTAAG	

ATAAATAAAT	TTGGCTTGCT	GTATTGGCCA	GGAAAAGAAA	GTATTAAAGT	ATGCATGCAT	1860
TATTTATTTA	AACCGAACGA	CATAACCGGT	CCTTTTCTTT	CATAATTTCA	TACGTACGTA	
GTGCACCAGG	GTGTTATTTA	ACAGAGGTAT	GTAACCTCTAT	AAAAGACTAT	AATTTACAGG	1920
CACGTGGTCC	CACAATAAAT	TGTCTCCATA	CATTGAGATA	TTTTCTGATA	TTAAATGTCC	
ACACGGAAAT	GTGCACATTT	GTTTACTTTT	TTTCTTCCTT	TTGCTTTGGG	CTTGTGATTT	1980
TGTGCTTTTA	CACGTGTAAA	CAAATGAAAA	AAAGAAGGAA	AACGAAACCC	GAACACTAAA	
TGGTTTTTGG	TGTGTTTATG	TCTGTATTTT	GGGGGGTGGG	TAGGTTTAAG	CCATTGCACA	2040
ACCAAAAACC	ACACAAATAC	AGACATAAAA	CCCCCACC	ATCCAAATTC	GGTAACGTGT	
TTCAAGTTGA	ACTAGATTAG	AGTAGACTAG	GCTCATTGGC	CTAGACATTA	TGATTTGAAT	2100
AAGTTCAACT	TGATCTAATC	TCATCTGATC	CGAGTAACCG	GATCTGTAAT	ACTAAACTTA	
TTGTGTTGTT	TAATGCTCCA	TCAAGATGTC	TAATAAAAGG	AATATGGTTG	TCAACAGAGA	2160
AACACAACAA	ATTACGAGGT	AGTTCTACAG	ATTATTTTCC	TTATACCAAC	AGTTGTCTCT	
CGACAACAAC	AACAAA					
GCTGTTGTTG	TTGTTT					

MVCGSPGGML LLRAGLLALA ALCLLRVPGA RAAACEPVRI PLCKSLPWNM TKMPNHLHHS	60
TQANAILAIE QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE	120
PILIKYRHSW PENLACEELP VYDRGVCISP EAIVTADGAD FPMDSNGNC RGASSERCKC	180
KPIRATQKTY FRNNYNYVIR AKVKEIKTKC HDVTAVVEVK EILKSSLVNI PRDTVNLYTS	240
SGCLCPPLNV NEEYIIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLSK	300
SDSSNSDSTQ SQKSGRNSNP RQARN.	

Figure 9. Deduced amino acid sequence of human FRZB-1 protein. SEQ ID NO:9.

Figure 10. Nucleotide sequence of the full-length human FRZB-1 cDNA. SEQ ID NO:10.
This sequence was assembled from public ESTs from the Genbank database
(accession numbers: H18848, R63748, W38677, W44760, H38379 and N71244).

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GGCGGAGCGG GCCTTTTGGC GTCCACTGCG CGGCTGCACC CTGCCCCATC TGCCGGGATC      60
CCGCCTCGCC CGGAAAACCG CAGGTGACGC GCCGACGTGG GACGGGGTAG ACGGCCCTAG

ATGGTCTGCG GCAGCCCGGG AGGGATGCTG CTGCTGCGGG CCGGGCTGCT TGCCCTGGCT      120
TACCAGACGC CGTCGGGCCC TCCCTACGAC GACGACGCCC GGCCCGACGA ACGGGACCGA

GCTCTCTGCC TGCTCCGGGT GCCCGGGGCT CGGGCTGCAG CCTGTGAGCC CGTCCGCATC      180
CGAGAGACGG ACGAGGCCCA CGGGCCCCGA GCCCGACGTC GGACACTCGG GCAGGCGTAG

CCCCTGTGCA AGTCCCTGCC CTGGAACATG ACTAAGATGC CCAACCACCT GCACCACAGC      240
GGGACACGTC TCAGGGACGG GACCTTGATC TGATTCTACG GGTGGGTGGA CGTGGTGTCTG

ACTCAGGCCA ACGCCATCCT GGCCATCGAG CAGTTCGAAG GTCTGCTGGG CACCCACTGC      300
TGAGTCCGGT TGCGGTAGGA CCGGTAGCTC GTCAAGCTTC CAGACGACCC GTGGGTGACG

AGCCCCGATC TGCTCTTCTT CCTCTGTGCC ATGTACGCGC CCATCTGCAC CATTGACTTC      360
TCGGGGCTAG ACGAGAAGAA GGAGACACGG TACATGCGCG GGTAGACGTG GTAAGTGAAG

CAGCACGAGC CCATCAAGCC CTGTAAGTCT GTGTGCGAGC GGGCCCGGCA GGGCTGTGAG      420
GTCGTGCTCG GGTAGTTCGG GACATTGAGA CACACGCTCG CCCGGGCCGT CCCGACACTC

CCCATACTCA TCAAGTACCG CCACTCGTGG CCGGAGAACC TGGCCTGCGA GGAGCTGCCA      480
GGGTATGAGT AGTTCATGGC GGTGAGCACC GGCCTCTTGG ACCGGACGCT CCTCGACGCT

GTGTACGACA GGGGCGTGTG CATCTCTCCC GAGGCCATCG TTAAGTGGGA CGGAGCTGAT      540
CACATGCTGT CCCCACACAC GTAGAGAGGG CTCCGGTAGC AATGACGCTT GCCTCGACTA

TTTCCTATGG ATTCTAGTAA CGGAACTGT AGAGGGGCAA GCAGTGAACG CTGTAAATGT      600
AAAGGATACC TAAGATCATT GCCTTTGACA TCTCCCCTGT CGTCACTTGC GACATTTACA

AAGCCTATTA GAGCTACACA GAAGACCTAT TTCCGGAACA ATTACAATA TGTCATTTCGG      660
TTCGGATAAT CTCGATGTGT CTTCTGGATA AAGGCCTTGT TAATGTTGAT ACAGTAAGCC

GCTAAAGTTA AAGAGATAAA GACTAAGTGC CATGATGTGA CTGCAGTAGT GGAGGTGAAG      720
CGATTTCATC TTCTCTATTT CTGATTCACG GTACTACACT GACGTCATCA CCTCCACTTC

GAGATTCTAA AGTCCTCTCT GTTAAACATT CCACGGGACA CTGTCAACCT CTATACCAGC      780
CTCTAAGATT TCAGGAGAGA CCATTTGTAA GGTGCCCTGT GACAGTTGGA GATATGGTCC

TCTGGCTGCC TCTGCCCTCC ACTTAATGTT AATGAGGAAT ATATCATCAT GGGCTATGAA      840
AGACCGACGG AGACGGGAGG TGAATTACAA TTAATCCTTA TATAGTAGTA CCCGATACTT

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GATGAGGAAC GTTCCAGATT ACTCTTGGTG GAAGGCTCTA TAGCTGAGAA GTGGAAGGAT 900
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 AGTGATTCTA GCAATAGTGA TTCCACTCAG AGTCAGAAGT CTGGCAGGAA CTCGAACCCC 1020
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 GTTTACTATA AAAATCATGT GATAACTGAT TATTACTTCT GTTCTCTTTT TGGTTTCTGC 1200
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 TTCTCTCTTC TCTCAACCCC TTTGTAATGG TTTGGGGGCA GACTCTTAAG TATATTGTGA 1260
 AAGAGAGAAG AGAGTTGGGG AAACATTACC AAACCCCGT CTGAGAATTTC ATATAACACT
 GTTTTCTATT TCACTAATCA TGAGAAAAAC TGTCTTTTGG CAATAATAAT AAATTAAACA 1320
 CAAAAGATAA AGTGATTAGT ACTCTTTTGG ACAAGAAAAC GTTATTATTA TTTAATTTGT
 TGCTGTTACC AGAGCCTCTT TGCTGAGTCT CCAGATGTTA ATTTACTTTC TGCACCCCAA 1380
 ACGACAATGG TCTCGGAGAA ACGACTCAGA GGTCTACAAT TAAATGAAAG ACGTGGGGTT
 TTGGGAATGC AATATTGGAT GAAAAGAGAG GTTCTTGGTA TTCACAGAAA GCTAGATATG 1440
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 TGTCACATAG GCAAAGCAAT CAAGCACCAG GAAGTGTTTA TGAGGAAACA ACACCCAAGA 1620
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 ACTTAATAAA AACTCTGACA GTCCTTCATT TTATTTATCC TCGAATTCTT TCTTGTAATA
 GCCTGATTGA GAAGCACAAC TGAAACCACT AGCCGCTGGG GTGTTAATGG TAGCATTCTT 1740
 CGGACTAACT CTTCTGTGTT ACTTTGGTCA TCGGCGACCC CACAATTACC ATCGTAAGAA
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 GGGTAACCAC TTTTCAAGTTT TTTTTCATTT TTT